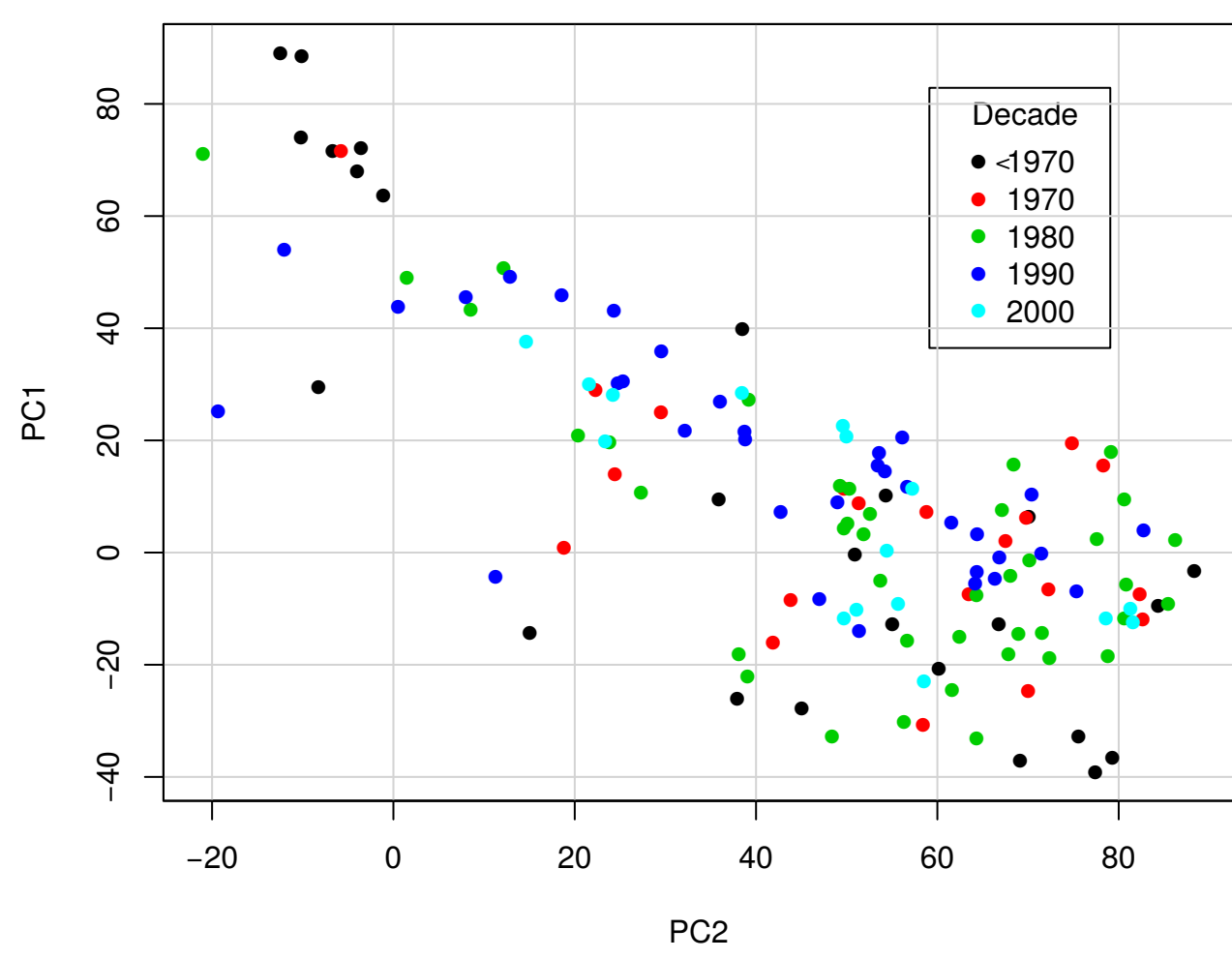
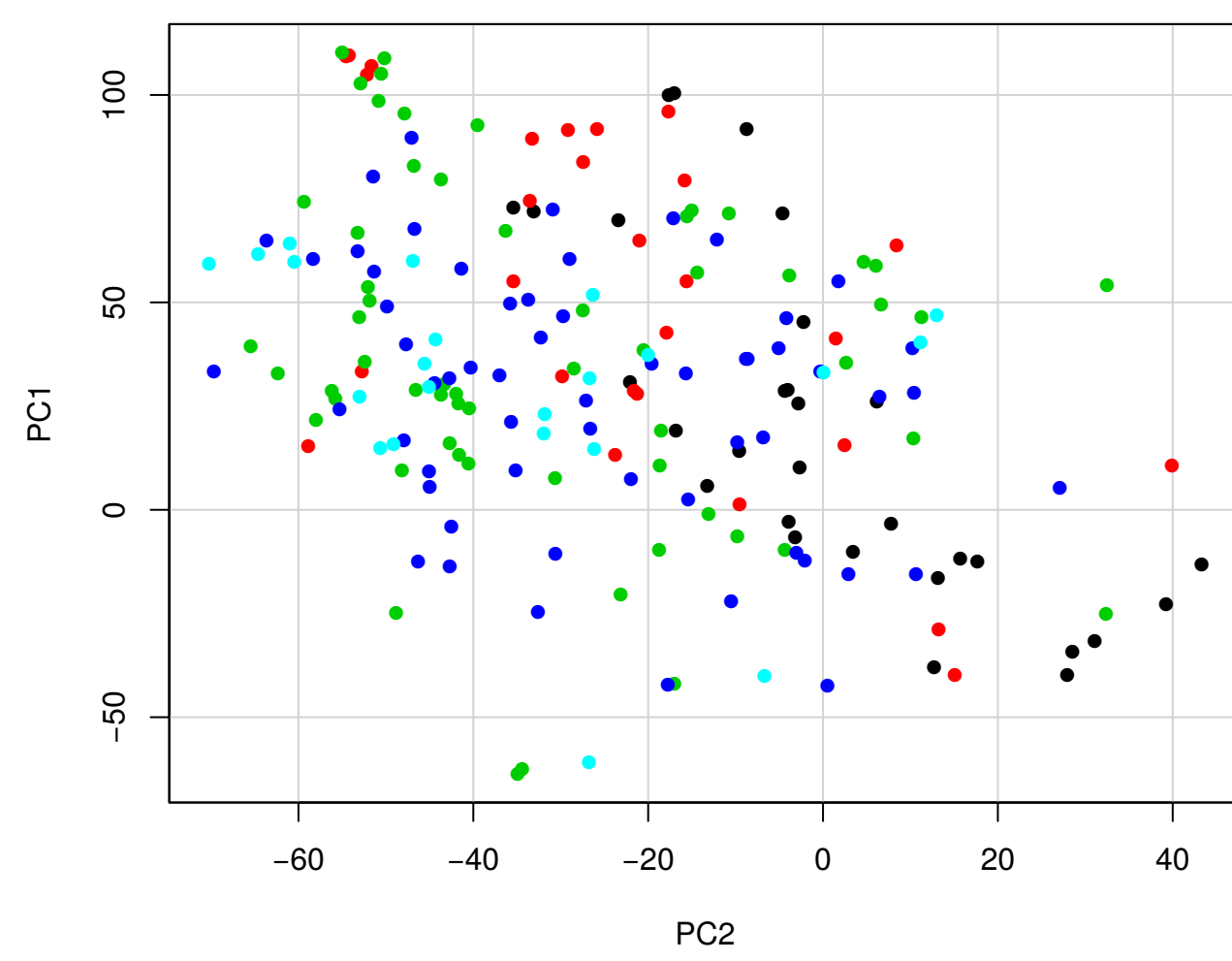


MG 0-I



MG III-IV



MG V+

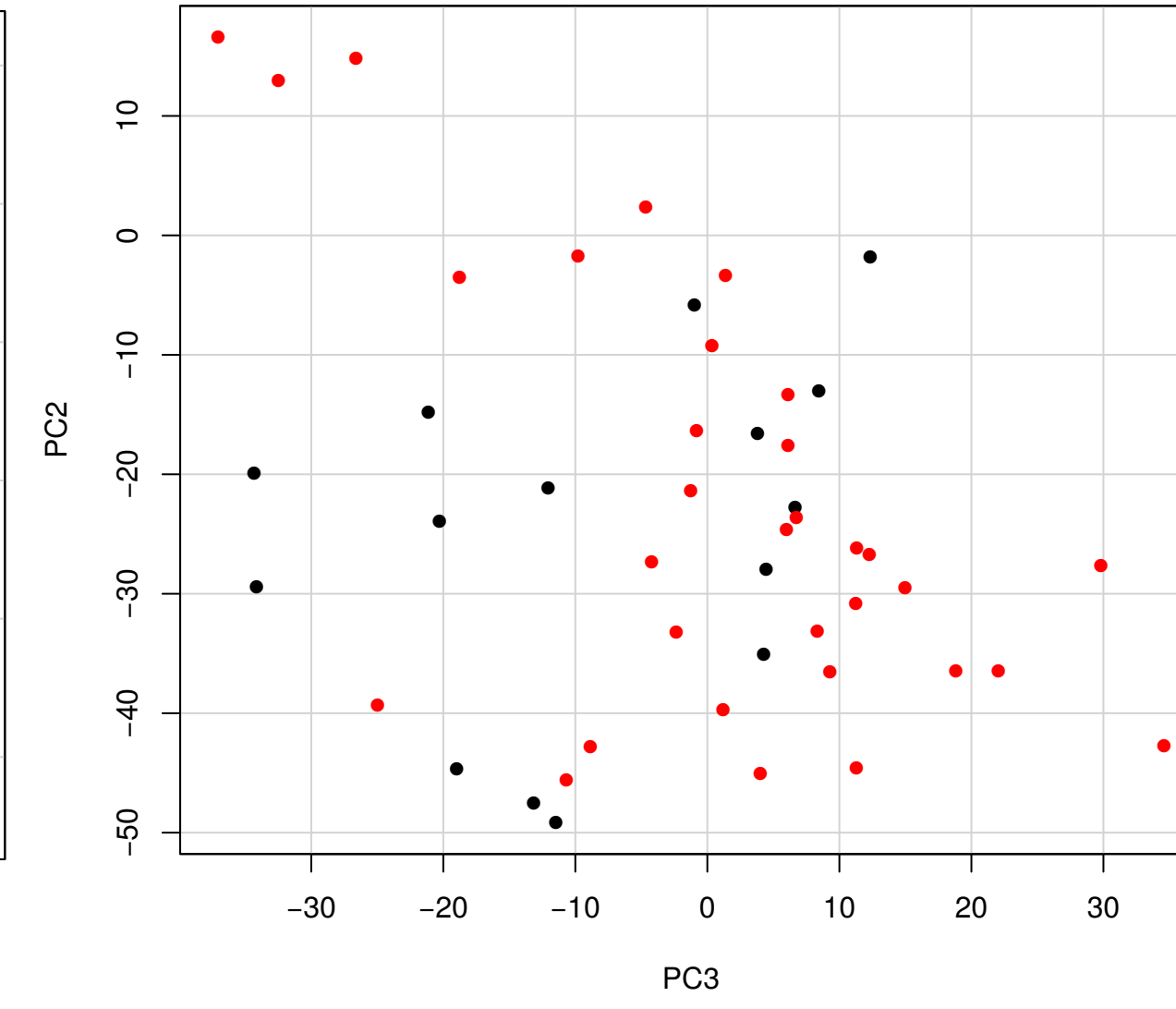
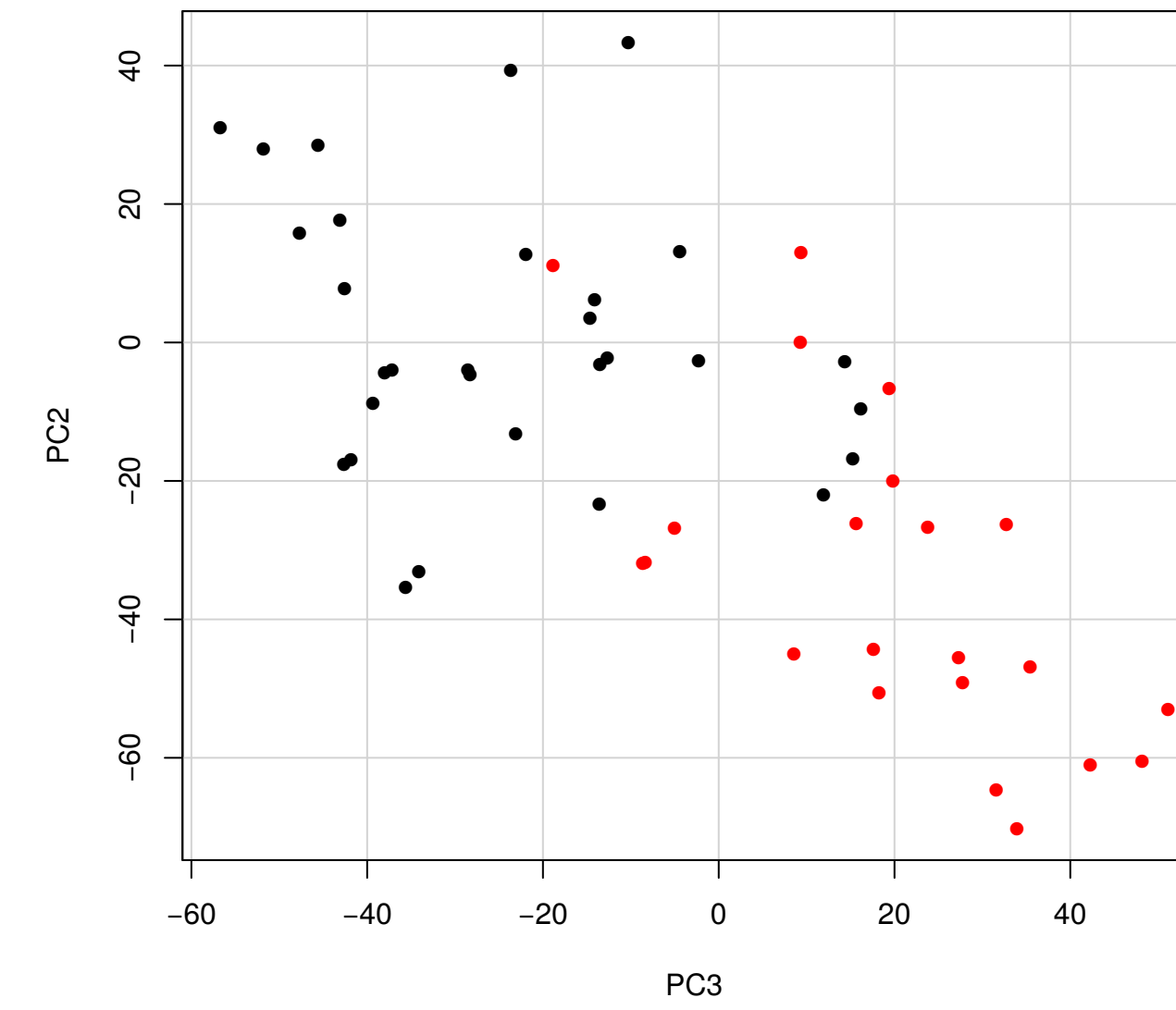
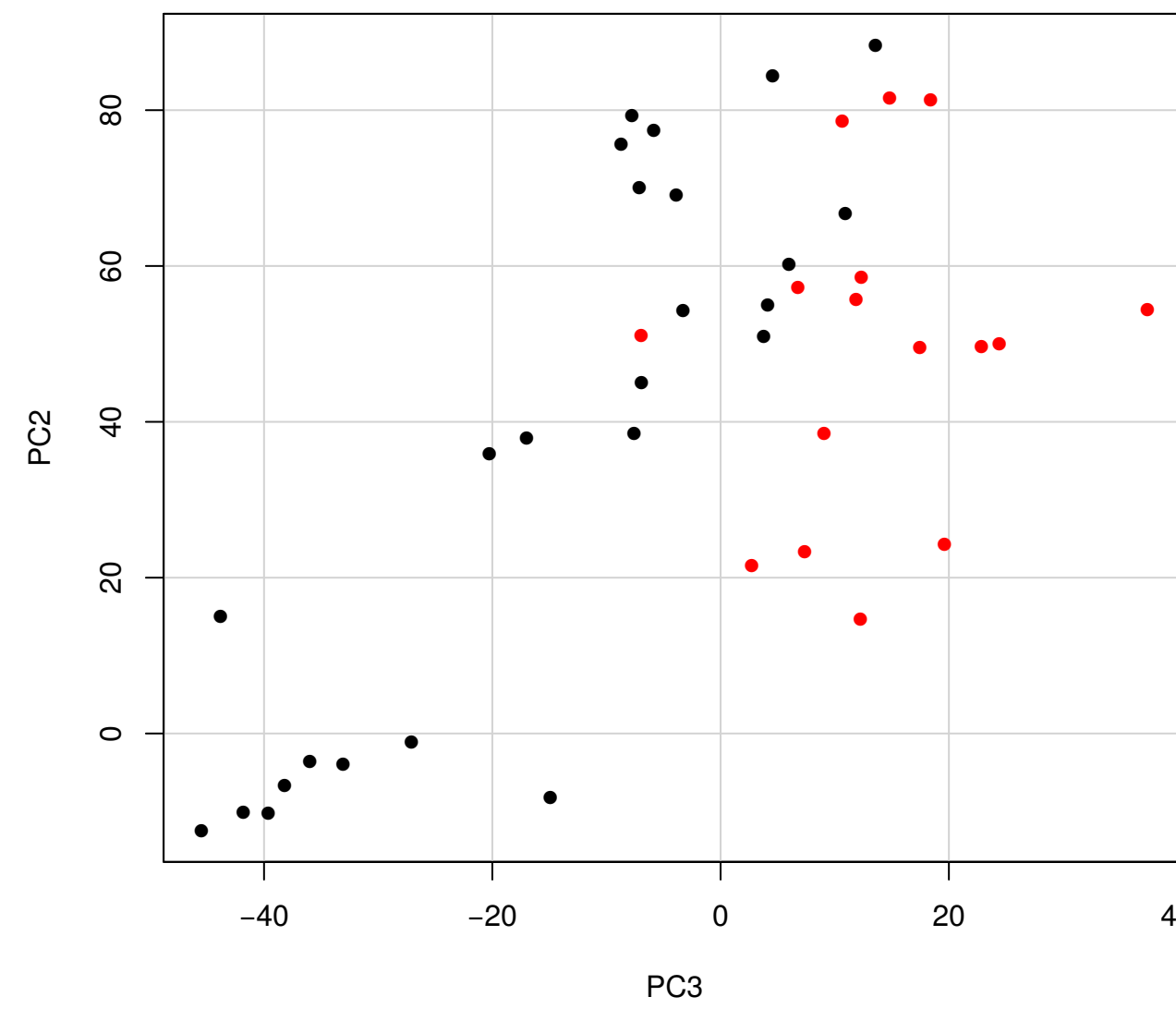
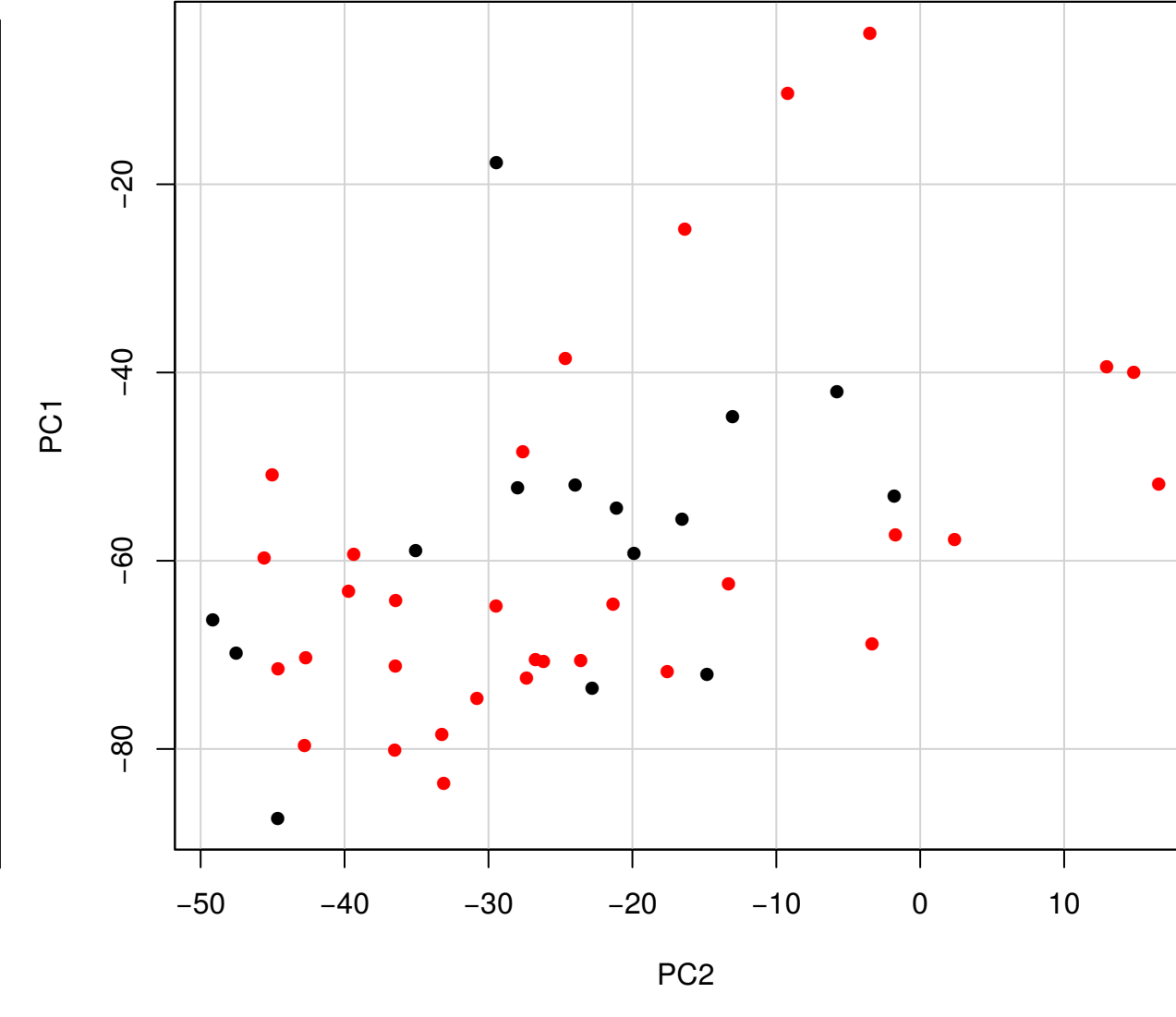
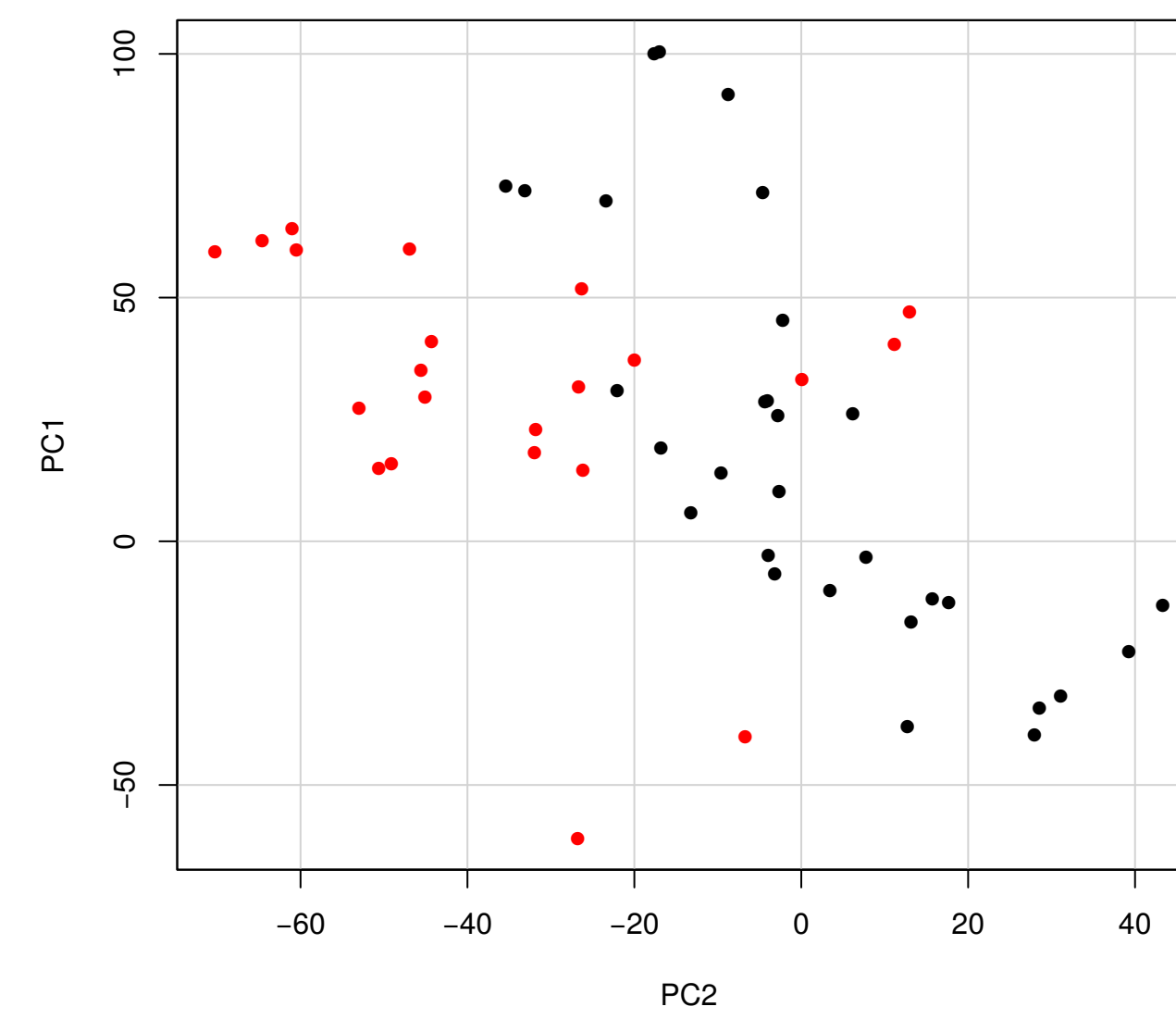
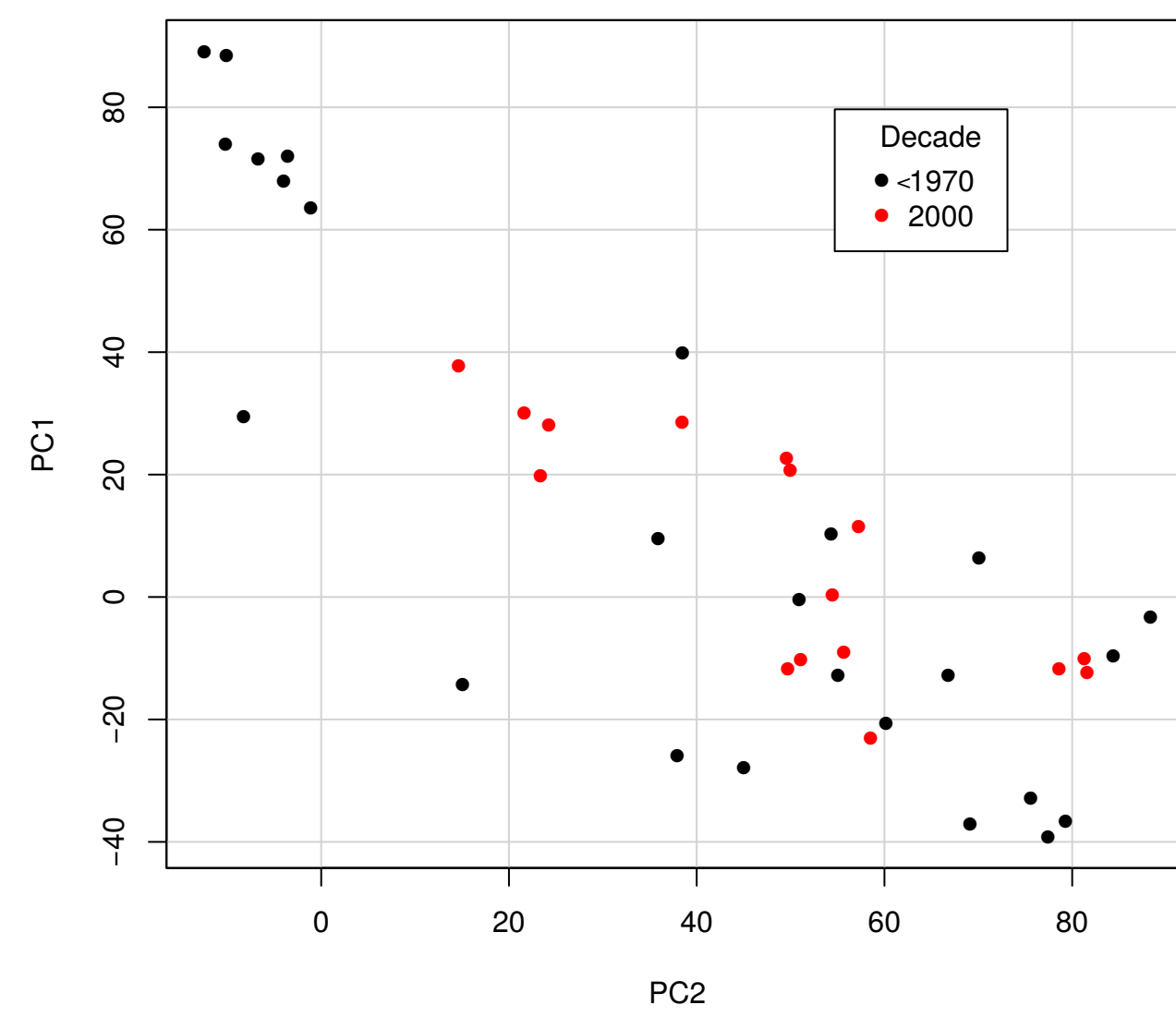
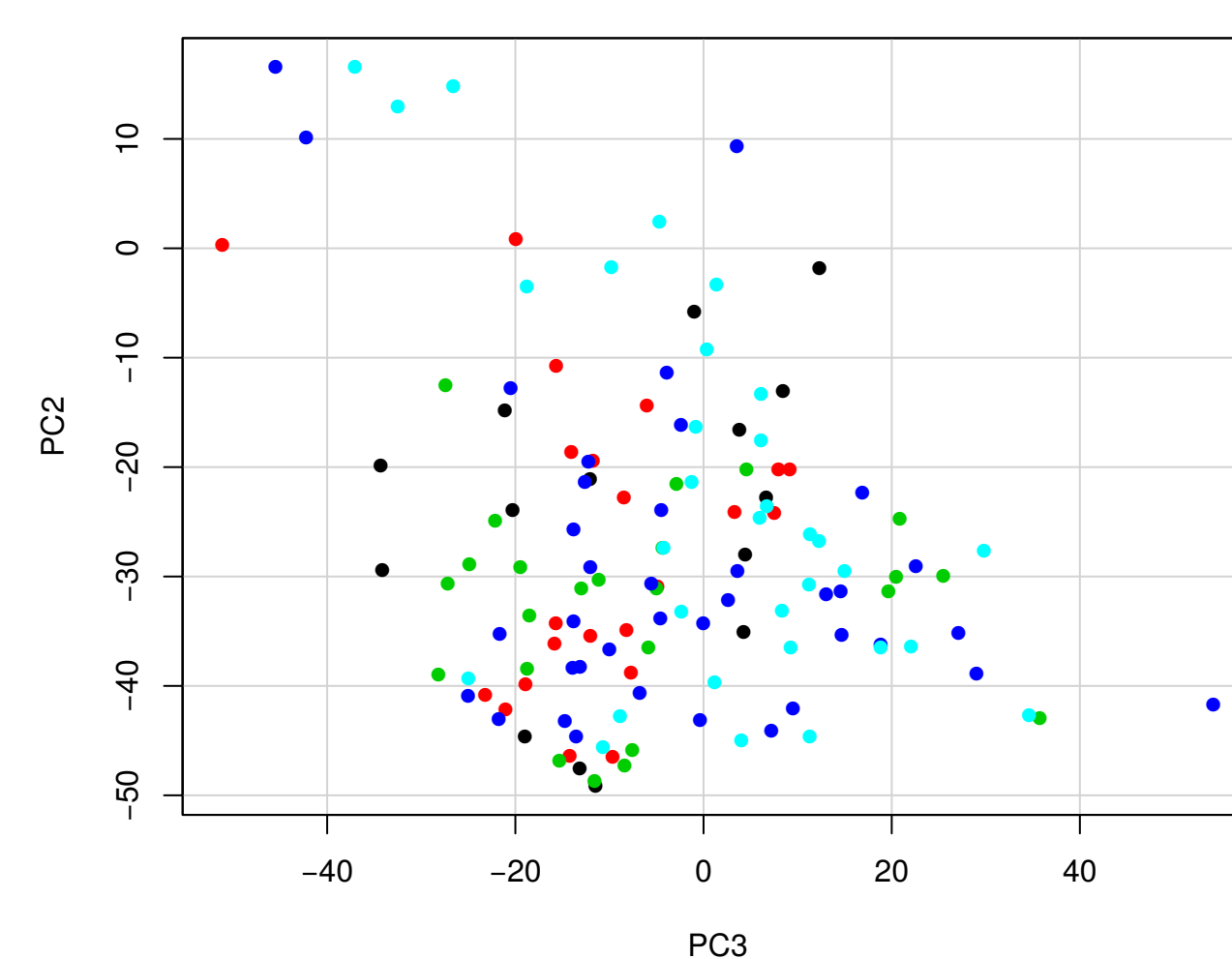
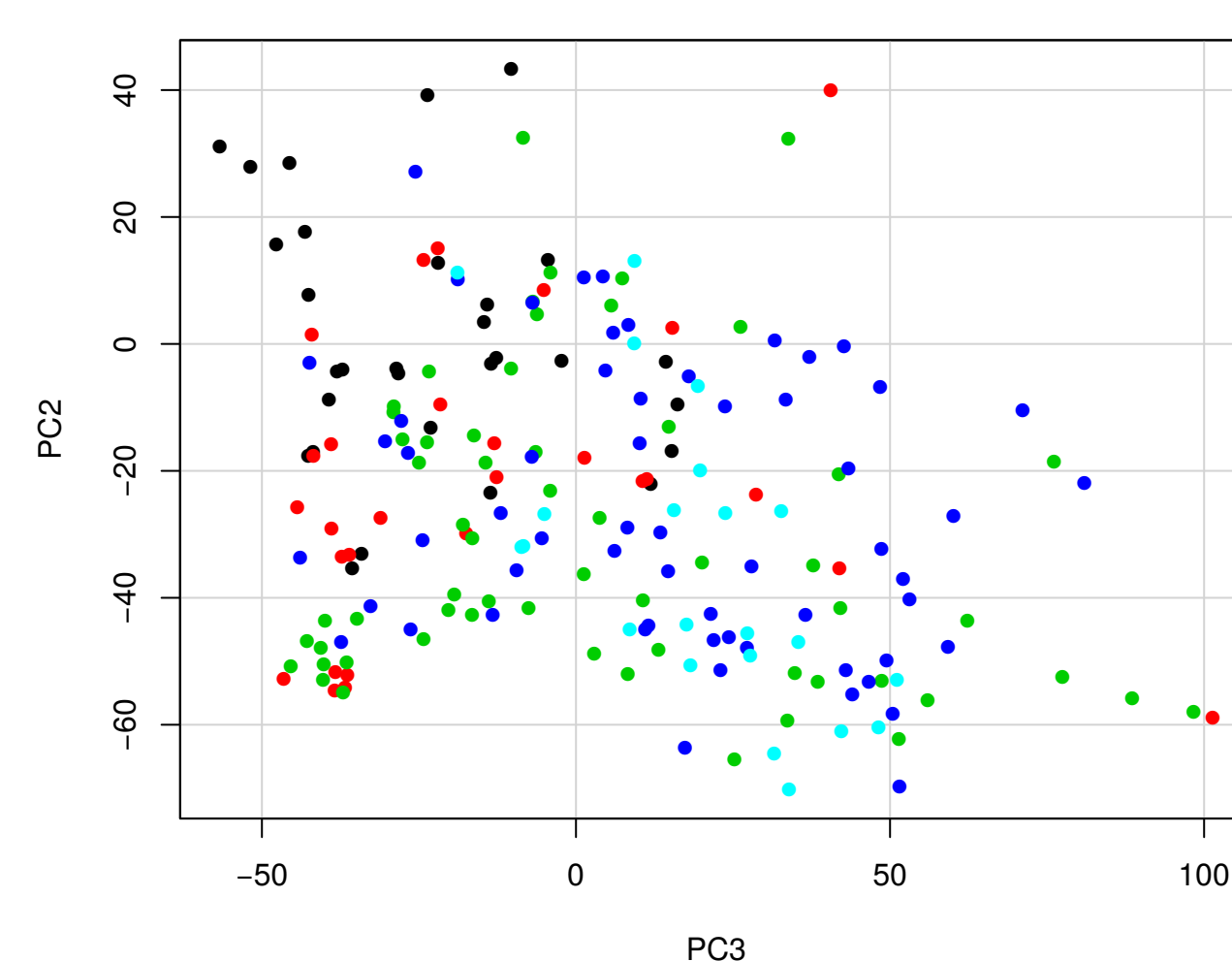
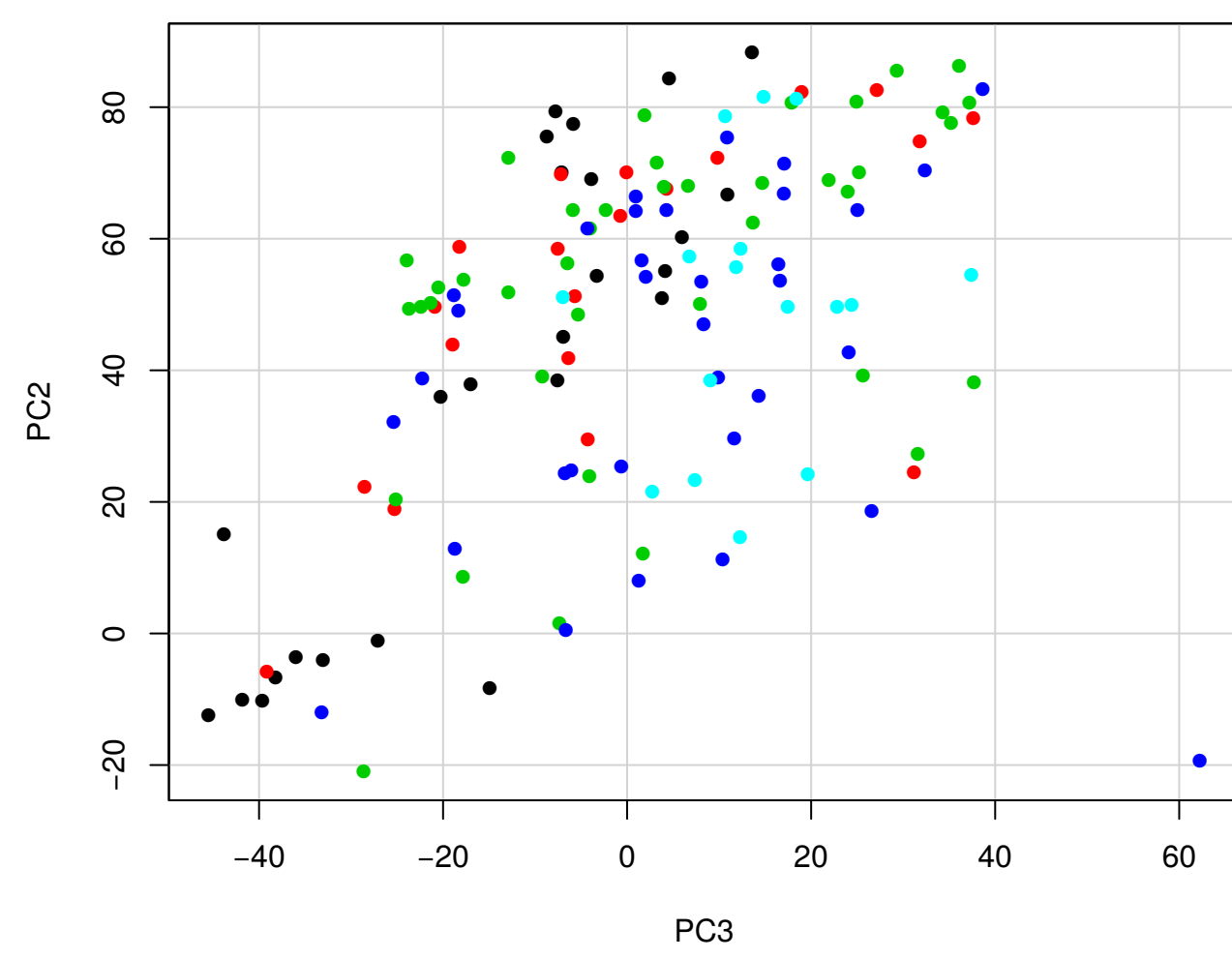
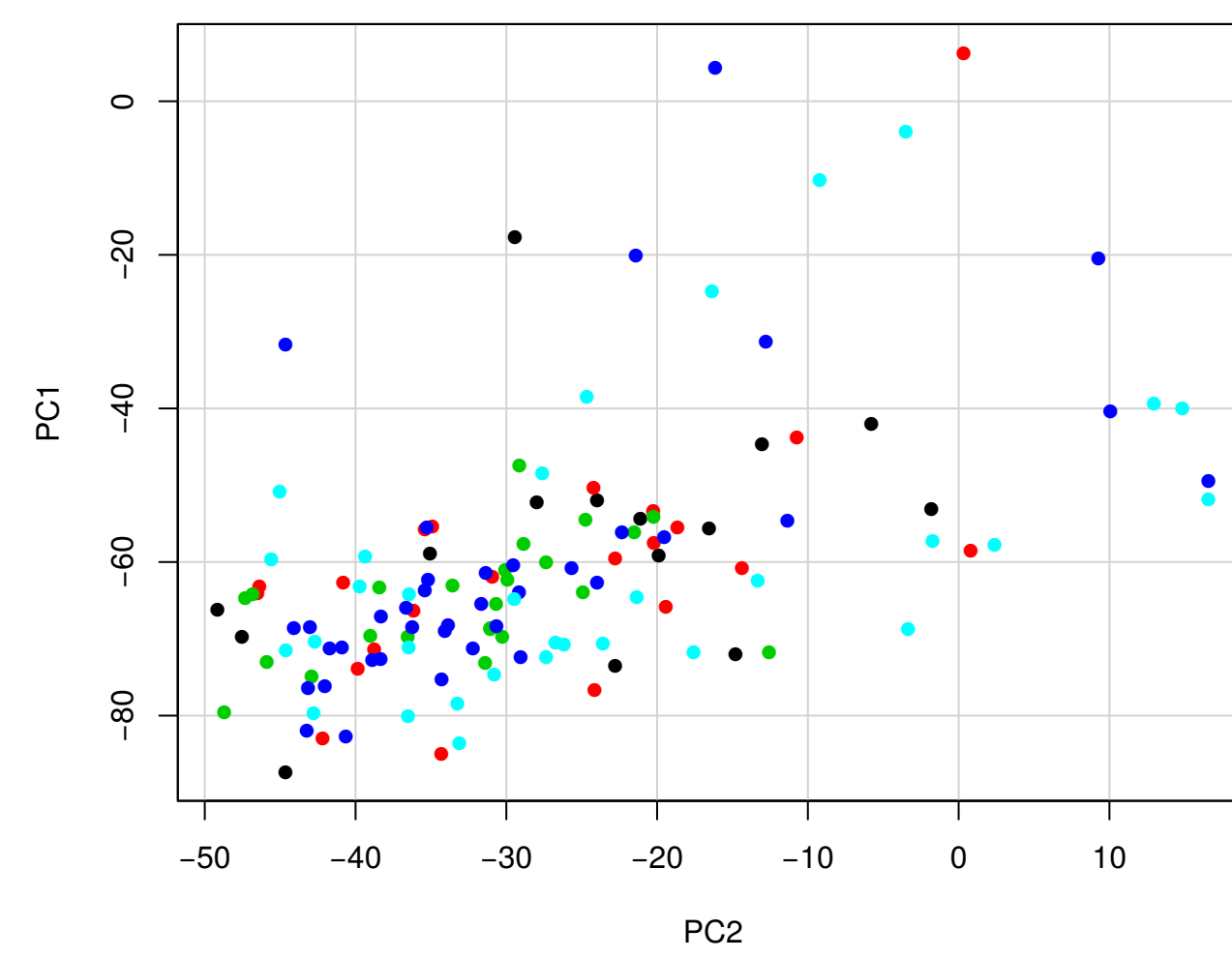


Figure S2: PC plots based on genome-wide marker information across all lines in the analysis. Lines are divided by population as discussed in main text. In the top two panels, lines are color coded based on the era of release. For clarity, only beginning and ending eras are shown in the bottom two panels.